

FIG._1A

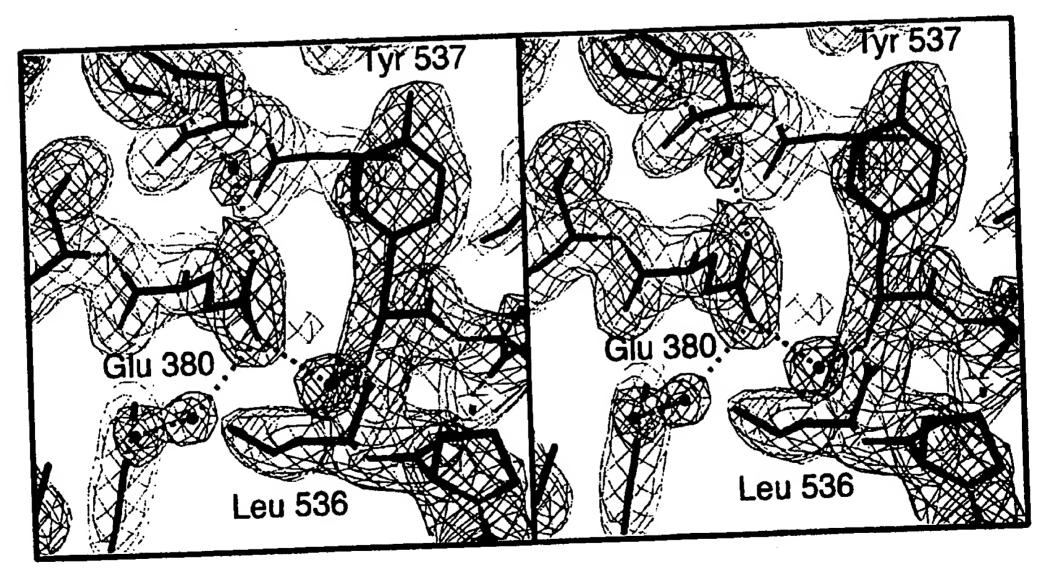
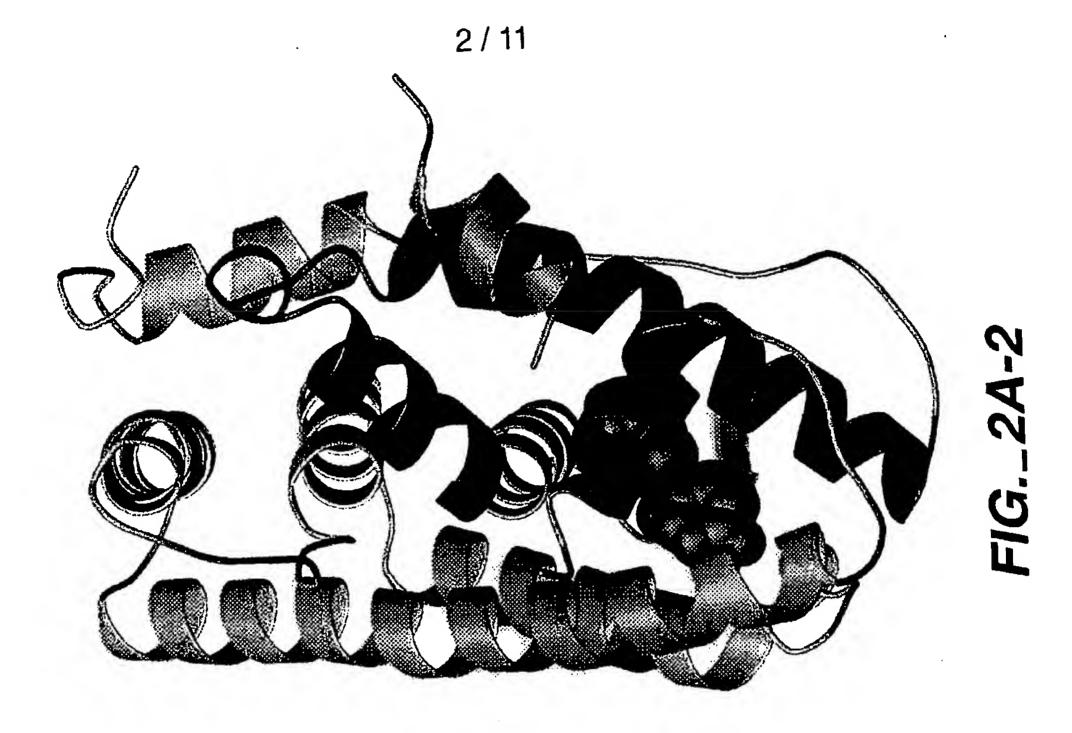
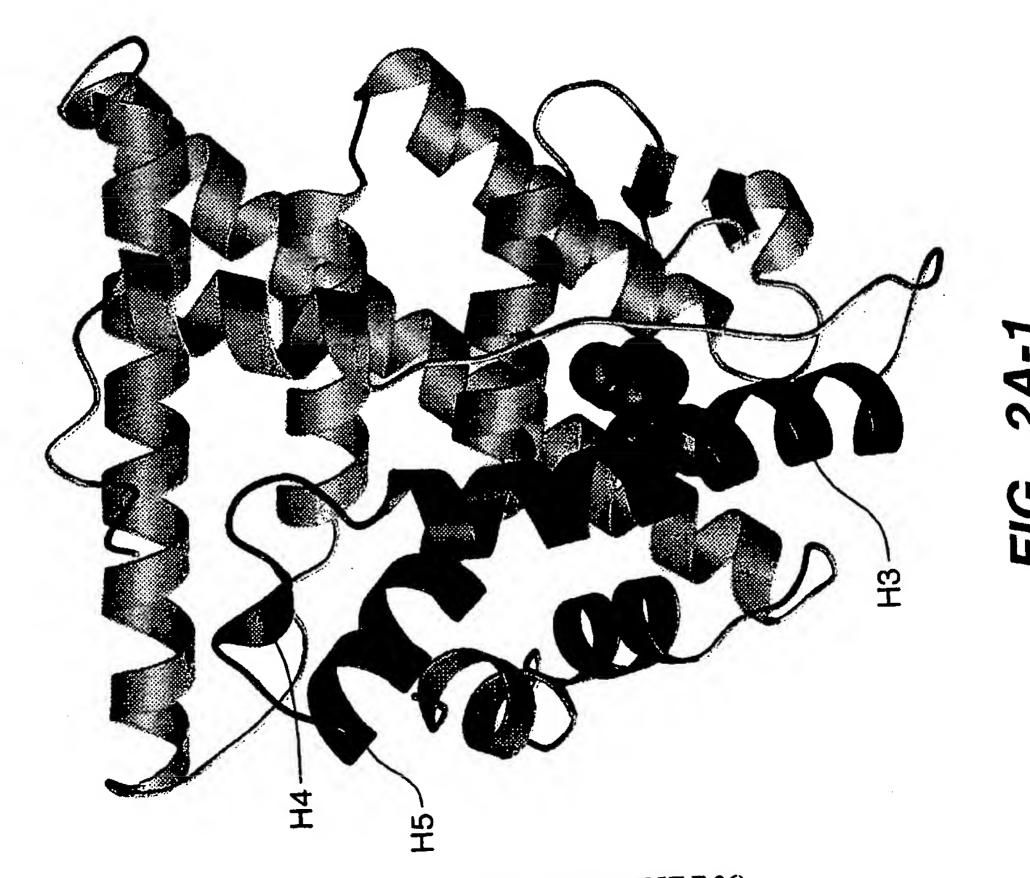


FIG._1B

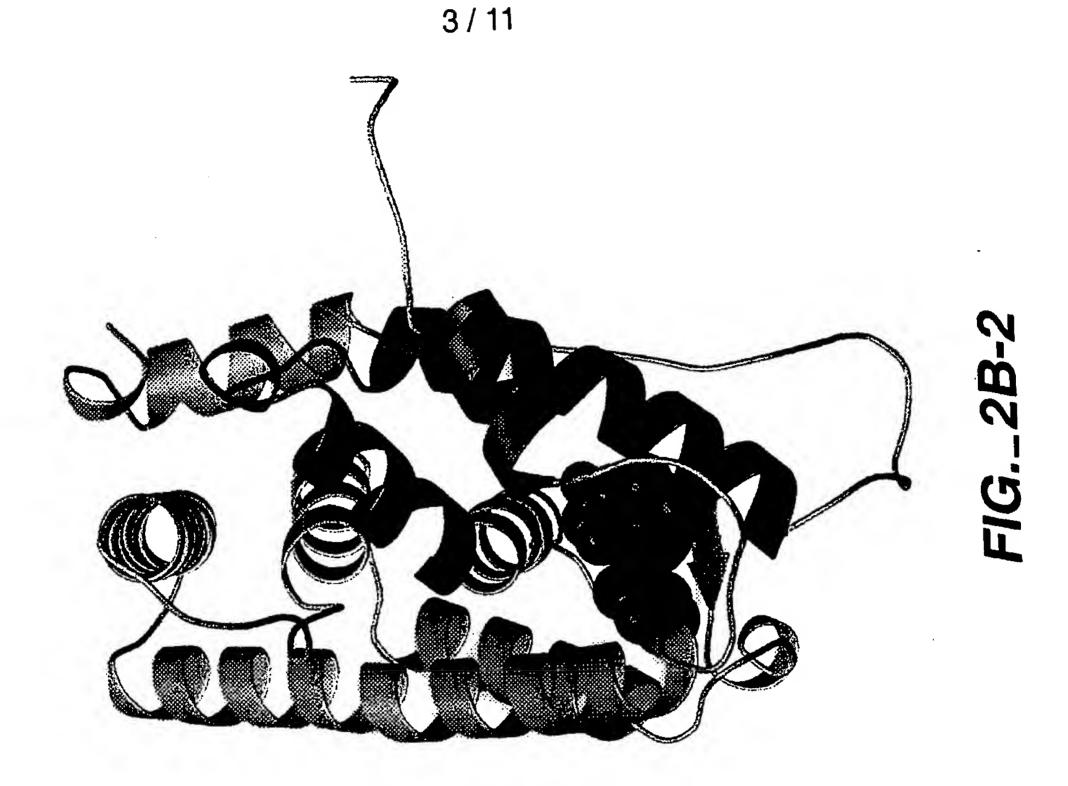
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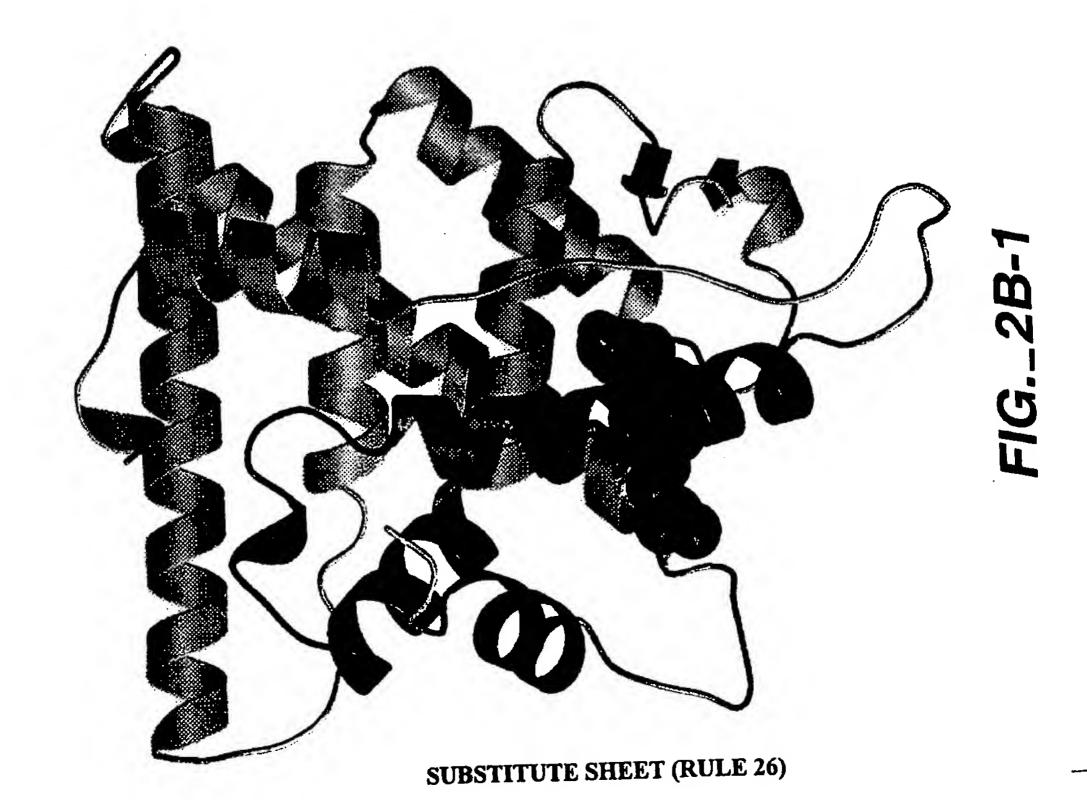




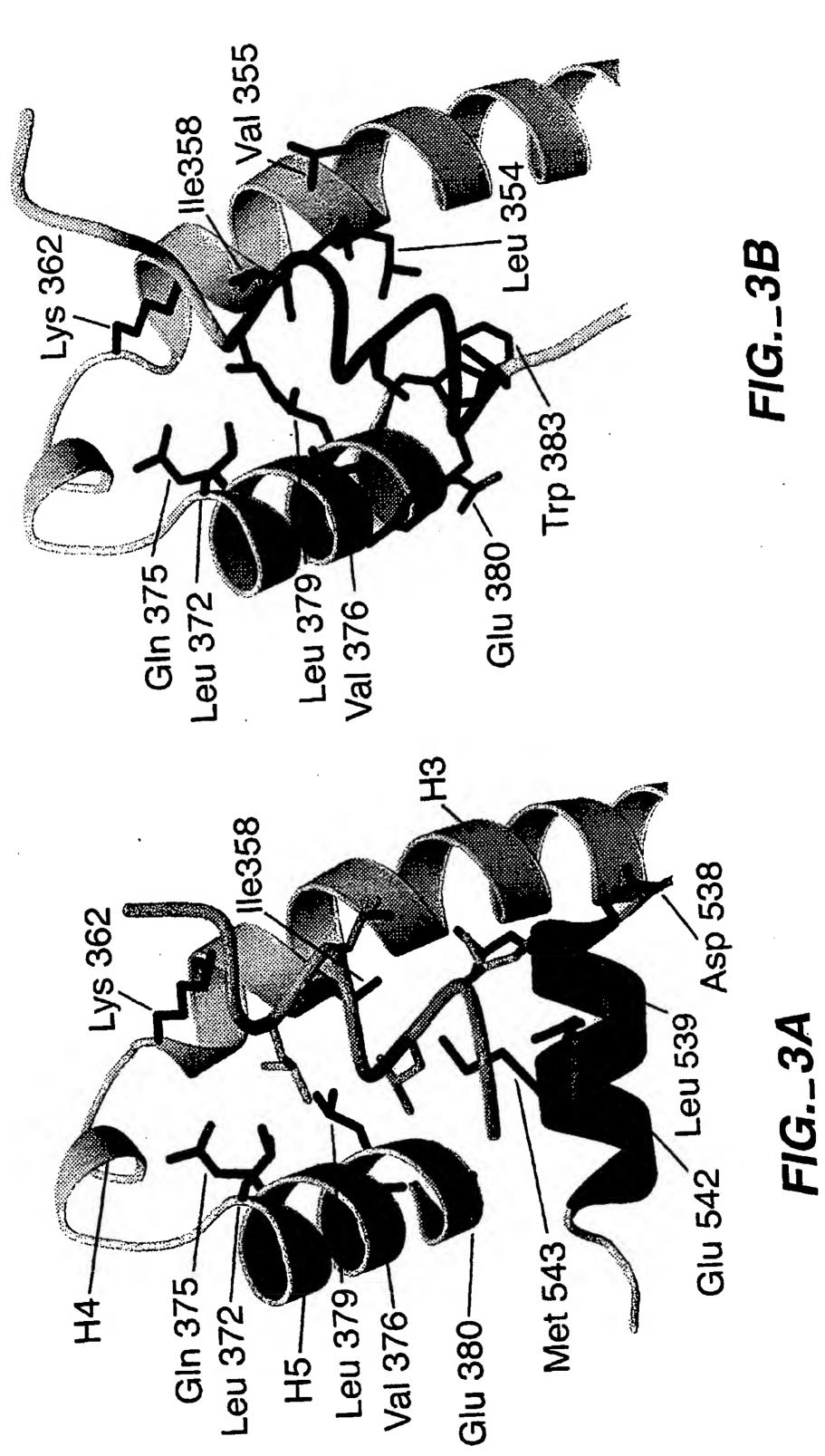
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CHRCTITHE CHEET (DITTE) (A)

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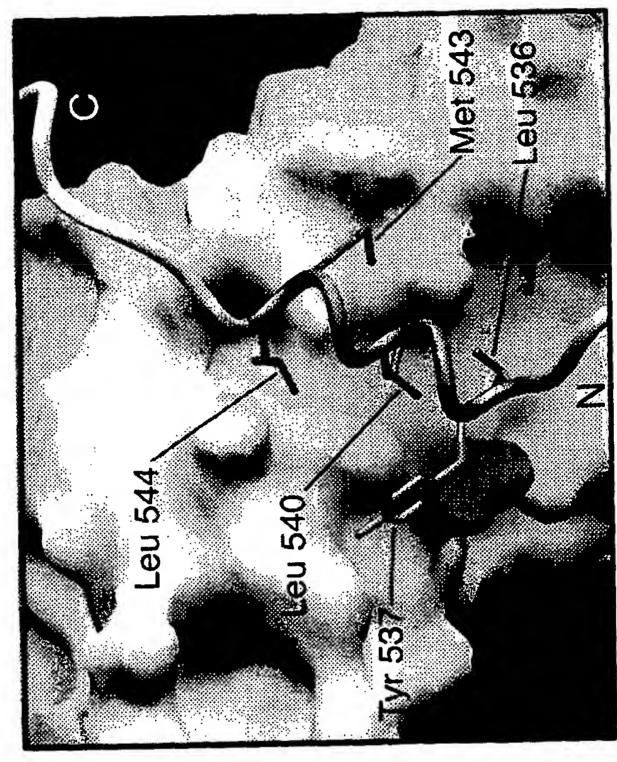


FIG._3D

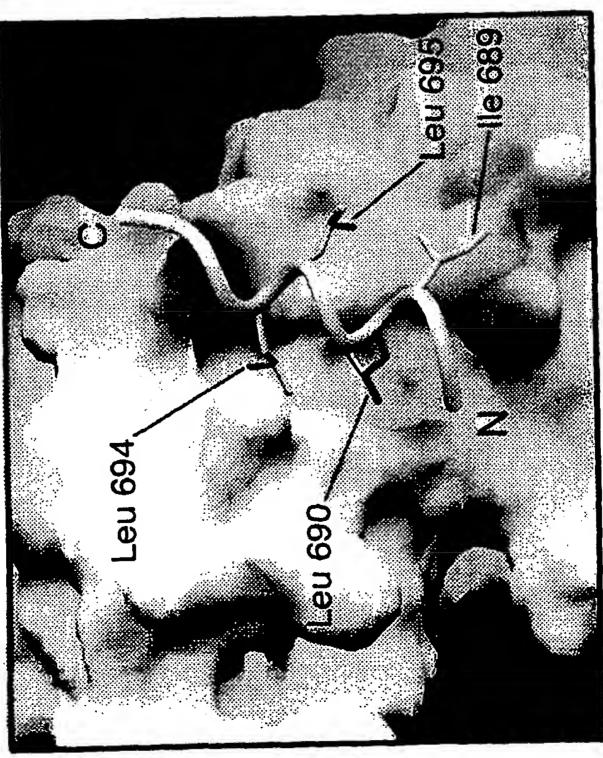
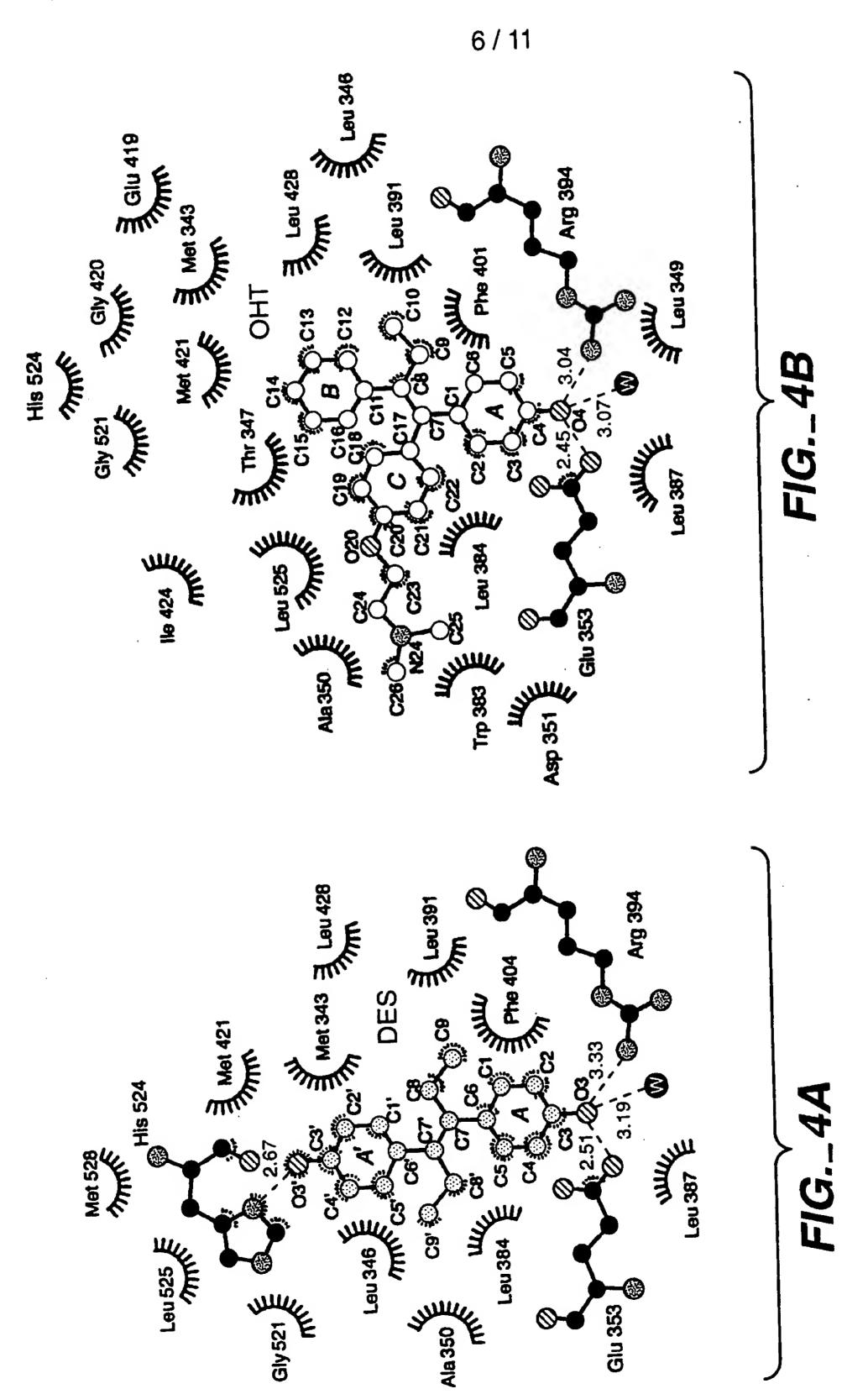


FIG._3C

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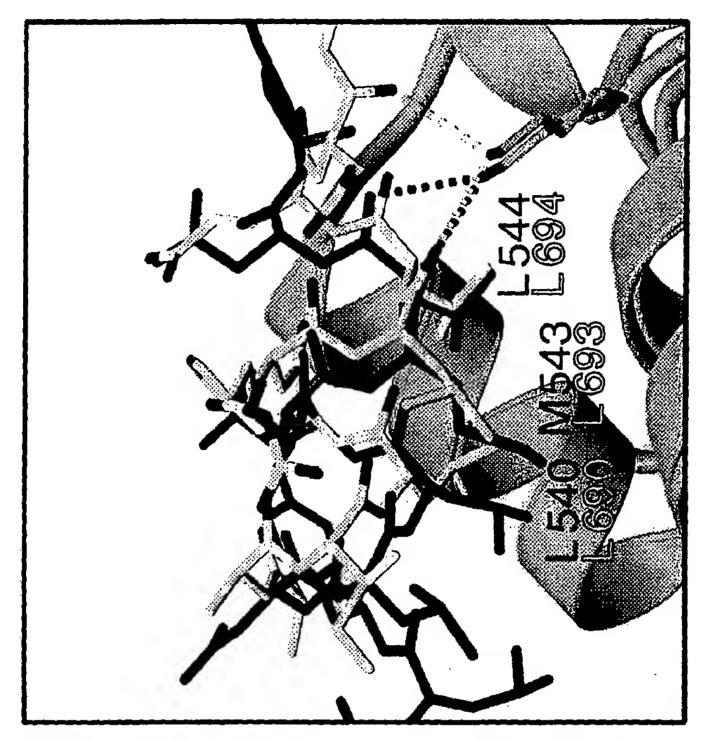
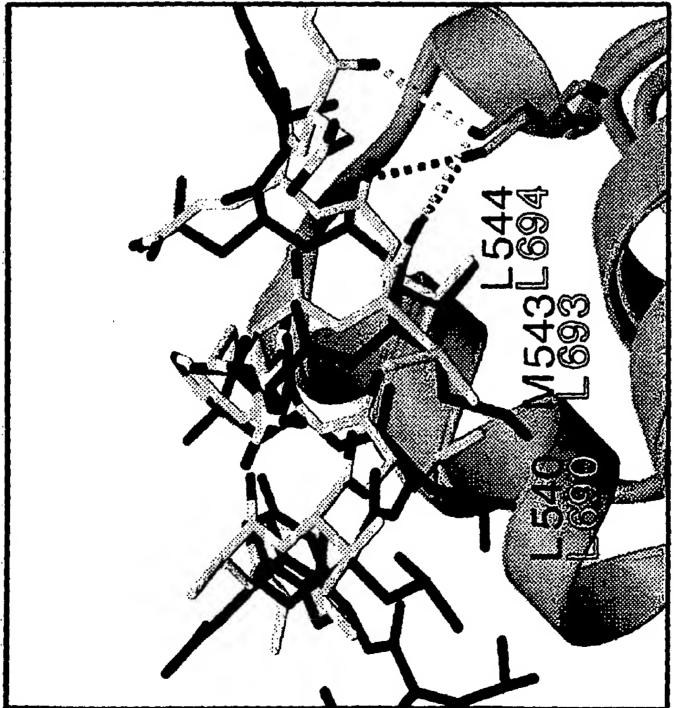


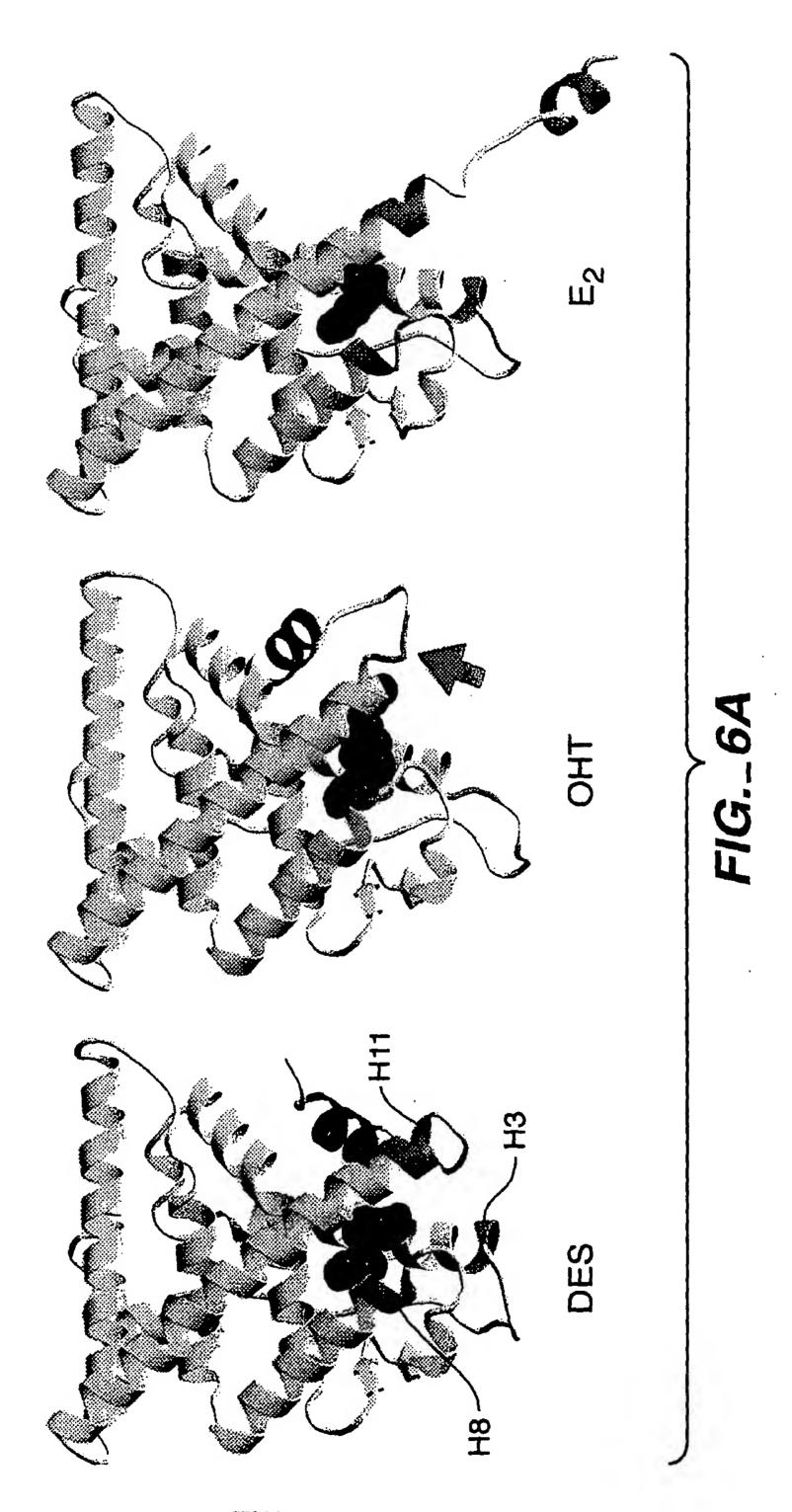
FIG._5B





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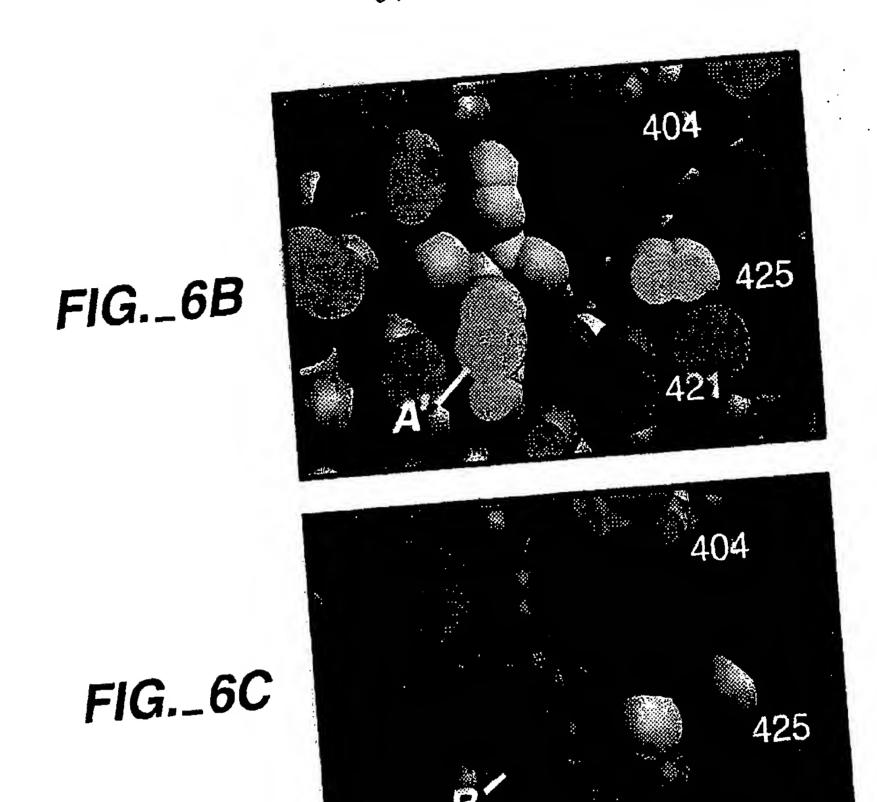
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SUBSTITUTE SHEET (RULE 26)

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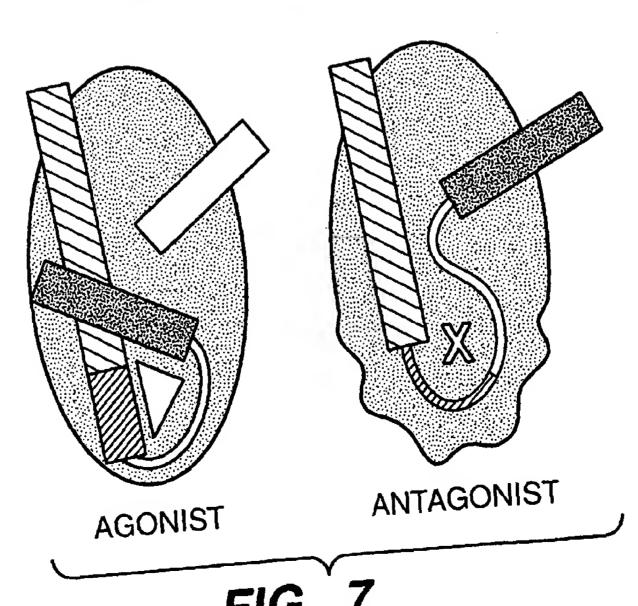
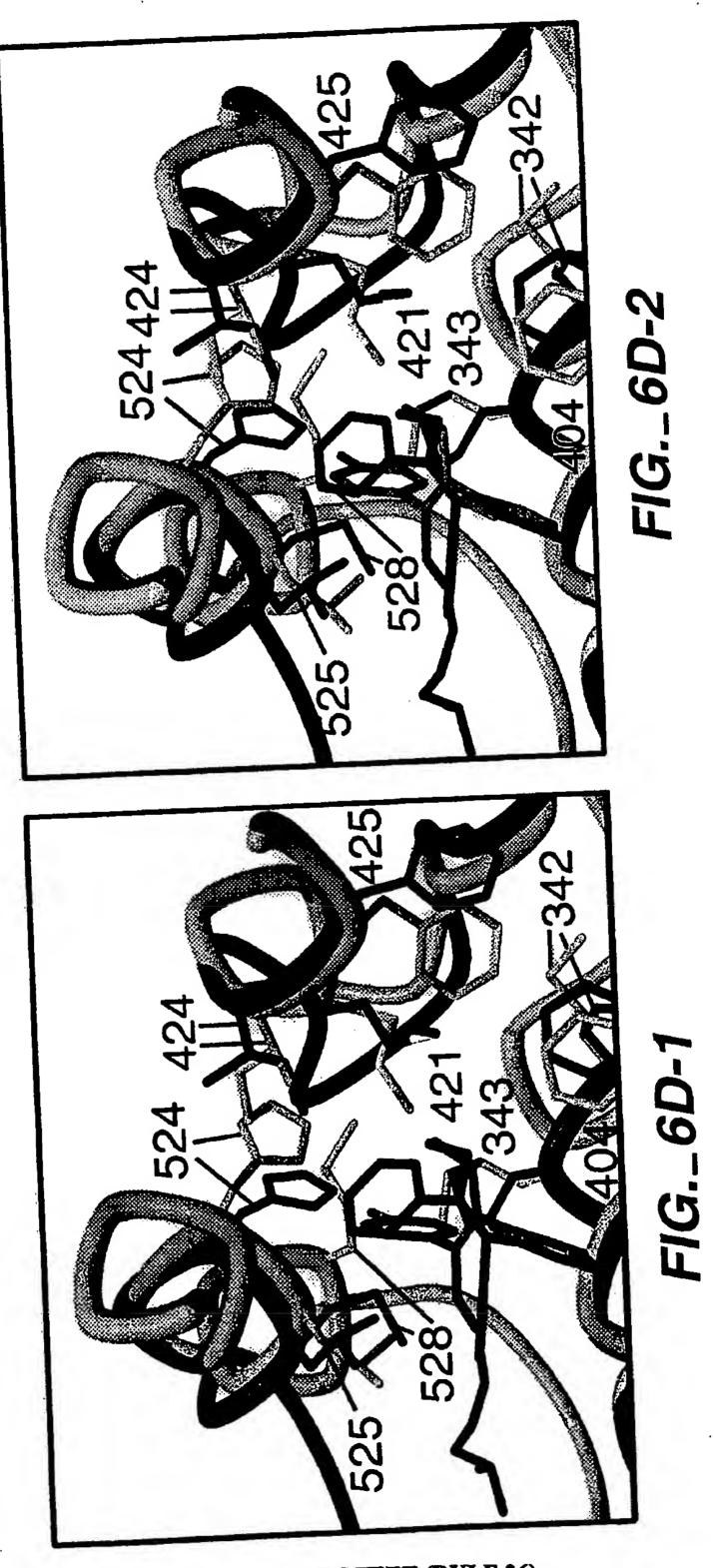


FIG._7



SUBSTITUTE SHEET (RULE 26)

hTRβ (SEQ ID NO:5) hRARγ (SEQ ID NO:7) hRXRα (SEQ ID NO:7) hPPARγ (SEQ ID NO:11) hPPARγ (SEQ ID NO:13) hVDR (SEQ ID NO:17) hFRα (SEQ ID NO:21) hMR (SEQ ID NO:23) hAR (SEQ ID NO:25)	461 (SEQ ID NO:6) 407 (SEQ ID NO:8) 418 (SEQ ID NO:10) 457 (SEQ ID NO:12) 473 (SEQ ID NO:14) 424 (SEQ ID NO:20) 546 (SEQ ID NO:20) 915 (SEQ ID NO:22) 966 (SEQ ID NO:24) 444 (SEQ ID NO:24)
309 GCC 309 h GCC 255 r AAC 267 h AGW 305 h YGV 320 h YGV 320 h SSA 267 h SSA 267 h YSW 600 h YSW 600 h YSW 600 h YSW 600 h	2 458 458 457459 LEVF REML REML REML REML REML REML REML REML
H5 302 306 306 301 305 •• CEDQIILLK CEDQIILLK LDDQVILLK LDDQVILLK SEDQIVLLK LDDQVILLE LDDQVILLE LDDQVILLE LDDQVILLE LDDQVILLE LDDQVILLE LDDQVILLE LDDQVILLE LDDQVITLLQ IDDQVITLLQ IDQVITLLQ IDQVITLQ IDQVITLQ IDQVITLLQ IDQVITLQ	H17 454 454 453 450 LFP 407 PMP 407 PMP 707 PMP 708 EFP 708 EFP 904 EFP 904 EFP 905 EFP 905 EFP 906 EFP 907 EMLA 908 EFP 908 EFP 908 EFP 908 EFP 909 EFP
H44 293 MFC MFC MFS GFT GGFR GGFR DLT GGFR NLH GGFR NLH NLH NLH NLH NLH NLH NLH NL	hTRB rTRC hRARy hRXRC hPR hGR hGR hAR
**************************************	8B 8B
H3 281 284 28 280 283 287 6 6 6 7 7 7 7 7 7 8 1 7 8 1 7 8 7 8 8 8 8 1 7 8 7 8	FIG.